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In the Claims:

Please cancel claim 17 without prejudice. Please amend claims 1-3, 6-8, 15-16, and 18-19 as shown below.

- 1. (Currently amended) An isolated protein <u>having phytase activity</u> comprising an amino acid sequence of SEQ ID NO:2 at its N-terminus wherein said protein having the following characteristics
 - (a) Molecular weight: about 47 kDa on SDS-PAGE.
 - (b) Optimal pH: pH 3.5 pH 4.5,
 - (c) Optimal temperature: 45°C 55°C.
 - (d) Substrate specificity: phytate, p-nitrophenyl phosphate, tetrasodium pyrophosphate, ATP or ADP,
 - (e) Michaelis constant of 0.3 0.5 mM utilizing phytate as a substrate,
 - (f) High resistance to protease such as pepsin, trypsin, papain, elastase or panereatin pancreatin,
 - (g) Specific activity to phytate: at least 1,500 units/mg amino acid residues 23-433 of SEO ID NO: 7.
- 2. (Currently amended) The protein as set forth in claim 1, wherein the protein comprises an amino acid sequence represented by amino acid residues 23 433 of SEQ ID-NO:7 or amino acid sequence having over 70% sequence homology with the same has the following characteristics:
 - (a) molecular weight: about 47 kDa on SDS-PAGE,
 - (b) optimal pH: pH 3.5-pH 4.5,
 - (c) Optimal temperature: 45 °C-55 °C,

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(d) substrate specificity: phytate, p-nitrophenyl phosphate, tetrasodium pyrophosphate, ATP or ADP,

- (e) michaelis constant of 0.3-0.5 mM utilizing phytate as a substrate,
- (f) high resistance to protease such as pepsin, trypsin, papain, elastase or pancreatin, and (g) specific activity to phytate; at least 1.500 units/mg.
- 3. (Currently amended) The protein as set forth in claim 1, wherein the protein comprises an amino acid a polypeptide sequence represented by of SEQ ID NO:7 or an amino acid sequence having over 70% sequence homology with the same.
- 4. (Canceled)
- (Previously presented) The protein as set forth in claim 1, wherein the specific activity
 of the protein to phytate is at least 3,000 units/mg
- 6. (Currently amended) A An isolated gene eoding encoding the protein of claim 1.
- 7. (Currently amended) The gene as set forth in claim 6, wherein the gene has a base polynucleotide sequence represented by of SEQ ID NO:6 or a base sequence having over 70% sequence homology with the same.
- 8. (Currently amended) \triangle An isolated microorganism belonging to Citrobacter species producing the protein of claim 1.

- (Previously presented) A feed additive containing the protein of claim 1 as an effective ingredient.
- (Original) The microorganism as set forth in claim 8, wherein Citrobacter species is Citrobacter braakii.
- 11. (Original) The microorganism as set forth in claim 10, wherein *Citrobacter braakii* is *Citrobacter braakii* YH-15 strain (Accession No: KCCM 10427).
- 12. (Previously presented) A feed additive containing the microorganism of claim 8 as an effective ingredient.
- 13. (Previously presented) The protein as set forth in claim 2, wherein the specific activity of the protein to phytate is at least 3,000 units/mg
- 14. (Previously presented) The protein as set forth in claim 3, wherein the specific activity of the protein to phytate is at least 3,000 units/mg
- 15. (Currently amended) A An isolated gene coding the protein of claim 2.
- 16. (Currently amended) A An isolated gene coding the protein of claim 3.
- 17. (Canceled)

- 18. (New) A <u>An isolated</u> microorganism belonging to *Citrobacter* species producing the protein of claim 2.
- 19. (New) A $\underline{An\ isolated}$ microorganism belonging to Citrobacter species producing the protein of claim 3.
- 20. (Previously presented) A feed additive containing the protein of claim 2 as an effective ingredient.